AMENDMENTS TO THE SPECIFICATION

<u>Please replace the paragraph beginning at page 5, line 5 with the following rewritten</u> paragraph:

FigureFIG. 2 shows the cDNA and deduced amino acid sequences of human PIN1 and homologies with other WW domain proteins and PPIases. Figure-FIG. 2A shows the Pin1 nucleotide sequence (SEQ ID NO: 1) and predicted Pin1-amino acid sequence (SEQ ID NO: 2) (isas indicated in one-letter code). The fusion points between GAL4 and Pin1 in six different isolated clones were: clone H20 at C-9; clone H16.24 and 38 at G:13G+13; clones H6 and H36 at C:15C+15. Underlined residues form a consensus bipartite nuclear localization signal. The N- and C-terminal boxes indicate the WW domain and PPIase domain_respectively. Nucleotide numbers are on the left and amino acid numbers on right. FigureFIG. 2B and 2C-shows alignments the alignment of the WW domain (B) and PPIase Domain (C) in selected proteins (from top to bottom SEQ ID NOs: 8-14). FIG. 2C shows the alignment of the PPIase domain in selected proteins (from top to bottom second p